SEQUENCE LISTING

<110> Cao, Liangxian
Trifillis, Panayiota

<120> METHODS FOR IDENTIFYING COMPOUNDS THAT MODULATE UNTRANSLATED REGION-DEPENDENT GENE EXPRESSION AND METHODS OF USING SAME

<130> 10589-012-999

<140> US 10/543,033
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<151> 2004-01-21

<150> 60/441,637
<151> 2003-01-21

<160> 90

<170> PatentIn version 3.2

<210> 1
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: consensus G-quartet element from synthetic sequences

<220>
<221> misc_feature
<222> 3, 7, 8, 11
<223> n = a, t, c, or g

<220>
<221> misc_feature
<222> (7)..(8)
<223> This represents one form of the sequence as described, other forms described may have up to five nucleotides in this variable region

<400> 1
ggntggnngg ntgg 14

<210> 2
<211> 14
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: synthetic G-quartet oligonucleotide

<220>
<221> misc_feature
<222> 3, 4, 7, 8, 11, 12
<223> n = a, t, g or c

<220>
<221> misc_feature
<222> 3, 4, 7, 8, 11, 12
<223> This represents one form of the sequence as described, other forms described have longer variable regions, typical is 2 - 10 nucleotides

<400> 2
ggnnggnngg nngg 14

<210> 3
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Antisense minus uORF NcoI primer

<400> 3
ggccccatgg ctccggctgg acccggctgg gacccggctg ggagggcgcg ggagggcgcg 60

g 61

<210> 4
<211> 19
<212> RNA
<213> Oryctolagus cuniculus

<220>
<223> subunit of 15-LOX-DICE

<400> 4
ccccrcccuc uuccccaag 19

<210> 5
<211> 152
<212> DNA
<213> Homo sapiens

<400> 5
gcagaggacc agctaagagg gagagaagca actacagacc ccccctgaaa acaaccctca 60

gacgccacat cccctgacaa gctgccaggc aggttctctt cctctcacat actgacccac 120

ggctccaccc tctctcccct ggaaaggaca cc 152

<210> 6
<211> 792
<212> DNA
<213> Homo sapiens

<400> 6
tgaggaggac gaacatccaa ccttcccaaa cgcctcccct gccccaatcc ctttattacc 60

ccctccttca gacaccctca acctcttctg gctcaaaaag agaattgggg gcttagggtc 120

ggaacccaag cttagaactt taagcaacaa gaccaccact tcgaaacctg ggattcagga 180

atgtgtggcc tgcacagtga attgctggca accactaaga attcaaactg gggcctccag 240

aactcactgg ggcctacagc tttgatccct gacatctgga atctggagac cagggagcct 300

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acccaatgta ggagctgcct tggctcagac atgttttccg tgaaaacgga gctgaacaat 600

aggctgttcc catgtagccc cctggcctct gtgccttctt ttgattatgt tttttaaaat 660

atttatctga ttaagttgtc taaacaatgc tgatttggtg accaactgtc actcattgct 720

gagcctctgc tccccagggg agttgtgtct gtaatcgccc tactattcag tggcgagaaa 780

taaagtttgc tt 792

<210> 7
<211> 21
<212> RNA
<213> Homo sapiens

<220>
<223> Group I AU-Rich element(ARE) cluster of 3'untranslated region

<400> 7
auuuauuuau uuauuuauuu a 21

<210> 8
<211> 40
<212> DNA
<213> Homo sapiens

<400> 8
kctggaggat gtggctgcag agcctgctgc tcttgggcac 40

<210> 9
<211> 289
<212> DNA
<213> Homo sapiens

<400> 9
gccggggagc tgctctctca tgaaacaaga gctagaaact caggatggtc atcttggagg 60

gaccaagggg tgggccacag ccatggtggg agtggcctgg acctgccctg ggccacactg 120

accctgatac aggcatggca gaagaatggg aatattttat actgacagaa atcagtaata 180

tttatatatt tatattttta aaatatttat ttatttattt atttaagttc atattccata 240

tttattcaag atgttttacc gtaataatta ttattaaaaa tatgcttct 289

<210> 10
<211> 7008
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Expression Vector pCMRI

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cgagcaaaat ttaagctaca acaaggcaag gcttgaccga caattgcatg aagaatctgc 180

ttagggttag gcgttttgcg ctgcttcgcg atgtacgggc cagatatacg cgttgacatt 240

gattattgac tagttattaa tagtaatcaa ttacggggtc attagttcat agcccatata 300

tggagttccg cgttacataa cttacggtaa atggcccgcc tggctgaccg cccaacgacc 360

cccgcccatt gacgtcaata atgacgtatg ttcccatagt aacgccaata gggactttcc 420

attgacgtca atgggtggag tatttacggt aaactgccca cttggcagta catcaagtgt 480

atcatatgcc aagtacgccc cctattgacg tcaatgacgg taaatggccc gcctggcatt 540

atgcccagta catgacctta tgggactttc ctacttggca gtacatctac gtattagtca 600

tcgctattac catggtgatg cggttttggc agtacatcaa tgggcgtgga tagcggtttg 660

actcacgggg atttccaagt ctccacccca ttgacgtcaa tgggagtttg ttttggcacc 720

aaaatcaacg ggactttcca aaatgtcgta acaactccgc cccattgacg caaatgggcg 780

gtaggcgtgt acggtgggag gtctatataa gcagagctct ctggctaact aagctttcgg 840

cgcgccgagg taccatggga tccgaagacg ccaaaaacat aaagaaaggc ccggcgccat 900

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gcgcgttatt tatcggagtt gcagttgcgc ccgcgaacga catttataat gaacgtgaat 1200

tgctcaacag tatgaacatt tcgcagccta ccgtagtgtt tgtttccaaa aaggggttgc 1260

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<211> 47
<212> DNA
<213> Homo sapiens

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<210> 12
<211> 307
<212> DNA
<213> Homo sapiens

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taaaactata aatatggatc ttttatgatt ctttttgtaa gccctagggg ctctaaaatg 180

gtttacctta tttatcccaa aaatatttat tattatgttg aatgttaaat atagtatcta 240

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aaaaaaa 307

<210> 13
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<213> Homo sapiens

<220>
<223> Group III AU-Rich element(ARE) cluster of 3'untranslated region

<220>
<221> misc_feature
<222> (1)..(15)
<223> n = a, u, g or c

<400> 13
nauuuauuua uuuan 15

<210> 14
<211> 62
<212> DNA
<213> Homo sapiens

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ttctgccctc gagcccaccg ggaacgaaag agaagctcta tctcgcctcc aggagcccag 60

ct 62

<210> 15
<211> 427

<212> DNA
<213> Homo sapiens

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tcttggaaag tgtaggctta cctcaaataa atggctaact tatacatatt tttaaagaaa 360

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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Expression Vector pCMR2

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ccaacgaccc ccgcccattg acgtcaataa tgacgtatgt tcccatagta acgccaatag 180

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<213> Homo sapiens

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<212> DNA
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<211> 547
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<213> Homo sapiens

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aagcctaccc ccgcgccgcg ccctgcccgc cgctgcg 97

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<211> 99

<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 1 for amplify VEGF 5'UTR

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<210> 70
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<212> DNA
<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Primer 3 for amplify VEGF 5'UTR

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<210> 72
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 4 for amplify VEGF 5'UTR

<400> 72
aaacccgggc agcaaggcaa ggctccaatg cac 33

<210> 73
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 5 for amplify VEGF 3'UTR

<400> 73
gccgggcagg aggaaggagc ctcccctcagg gtttcggga 39

<210> 74
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer 6 for amplify VEGF 3'UTR

<400> 74
ctgcactaga gacaaagacg tgatgttaat 30

<210> 75
<211> 66
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Polylinker

<400> 75
gaacaaatgt cgacgggggc ccctagcaga tctagcgctg gatcccccgg ggagctcaug 60

gaagac 66

<210> 76
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for luciferase amplification

<400> 76
cggtgttggg cgcgttattt atcggagttg 30

<210> 77
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for luciferase amplification

<400> 77
ttggcgaaga atgaaaatag ggttggtact 30

<210> 78
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for GAPDH amplification

<400> 78
ggtgaaggtc ggagtcaacg ga 22

<210> 79
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer for GAPDH amplification

<400> 79
gagggatctc gctcctggaa g 21

<210> 80
<211> 55
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 5'UTR forward oligo

<400> 80

aaagtcgacg taaccgccag atttgaatcg cgggacccgt tggcagaggt ggcgg 55

<210> 81
<211> 54
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 5'UTR reverse oligo

<400> 81
aaaggatccg ggcaacgtcg gggcacccat gccgccgccg ccacctctgc caac 54

<210> 82
<211> 40
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 3'UTR forward oligo

<400> 82
aaagcggccg cggcctctgc cggagctgcc tggtcccaga 40

<210> 83
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 3'UTR reverse oligo

<400> 83
aaatctagac tcaggaacag ccgagatgac ctccaga 37

<210> 84
<211> 67
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SL top oligonucleotide

<400> 84
ctagaagctt agggccgcgg atccgcgcgc ggttcgccgc gcgcggatcc gcggtagcaa 60

gttagtc 67

<210> 85
<211> 68
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: SL bottom oligonucleotide

<400> 85
gactaagctt gctaccgcgg atccgcgcgc ggcgaaccgc gcgcggatcc gcggccctaa 60

gcttctag 68

<210> 86
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Sense/HindIII)

<400> 86
caagaagctt gcgcccggcc ccccaccccc cg 32

<210> 87
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Antisense/NcoI)

<400> 87
agcccatggt gctcactgcg gctccggccc c 31

<210> 88
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Sense/BglII)

<400> 88
agactctgaa ccagaaggcc aa 22

<210> 89
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PCR primer (Antisense/KpnI)

<400> 89
ctcggtacca gttttccaaa atatatttgc aaatgg 36

<210> 90
<211> 58
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: sense minus uORF HindIII primer

<400> 90
cccaagcttc gcgcccggcc ccccaccccct cgcagcaccc cgcgccccgc gccctccc 58